

Figure 1A

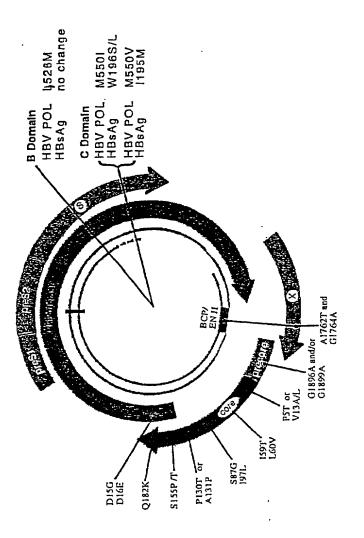


Figure 1B

(421)	430	,	440	450
422 SNDLSWLSLD	VSAAF	438 YHI _P PL	HPAAMPHLLIV	GSSGL ^D SRYVA
Do	main A			

HBSAg G112R	T123P	Y/F134	IS D144E G145R
460	470	480	490
464 466	477	488	499
RLSST _N SR _N NI*N	$N_{YH}Q_{XH}G^{X} \star \star \star D^{N}\Gamma H$	D _N Y _S CSRD _Q LYVS	$L_{M}LLY_{M}^{Q}T_{F}G_{M}^{R}$

HBsAq	A157D	E164.	D F170L	
500	510		520	530
	512	519	523/524/526/528/	′530 [°]
KLHLY _L S _A HPII _V	LGFRKILP	MG ^v G	GLSPFLLAQF	$\mathbf{T} SAIc_L s_A v_M v_T s_C R$
-	Domai	_		-

	W196L	W1995				
HBSAq	M195I/S196W M	1981 S204T	S210R			
540	550	560				
546	550 553	559	565			
AFF _P HCL _V A _V FS	Y MDDVLMVI	GAK _R s _T V _G Q _E H	$L_{SR}E_{F}L_{F}T_{A}S_{A}$			
Domain C						
570	580	590				
575	•					
IVTCNSFVLLSDL	GI HLNPNQF	KTKRW GYSLN	IFMGYI _V I G			
	Domain D		Oomain E			

Figure 2

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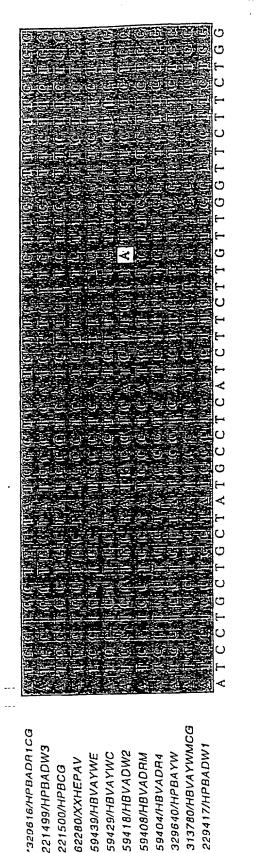
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Lie

P 108

329616/HPBADR1CG

221499/HPBADW3



221500/HPBCG

313780/HBVAYWMCG

329640/HPBAYW

59418/HBVADW2

59408/HBVADRM 59404/HBVADR4

59439/HBVAYWE 59429/HBVAYWC

62280/XXHEPAV

221500/HPBCG

229417/HPBADW1

H. H. H. Herbert Hand Bern Hand

Figure 3 continued

AAGGAAC

313780/HBVAYWMCG

329640/HPBAYW

59408/HBVADRM 59404/HBVADR4

59429/HBVAYWC 59418/HBVADW2

59439/HBVAYWE

229417/HPBADW1

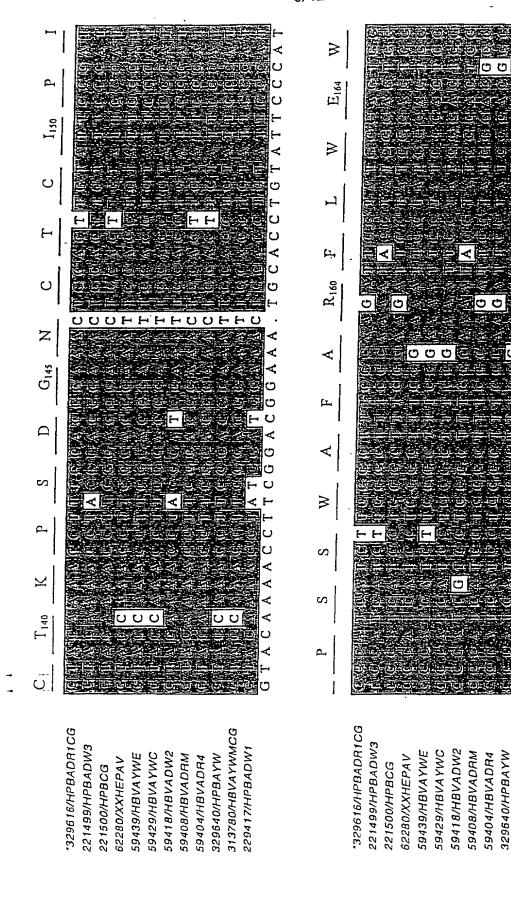


Figure 3 continued

313780/HBVAYWMCG

229417/HPBADW1

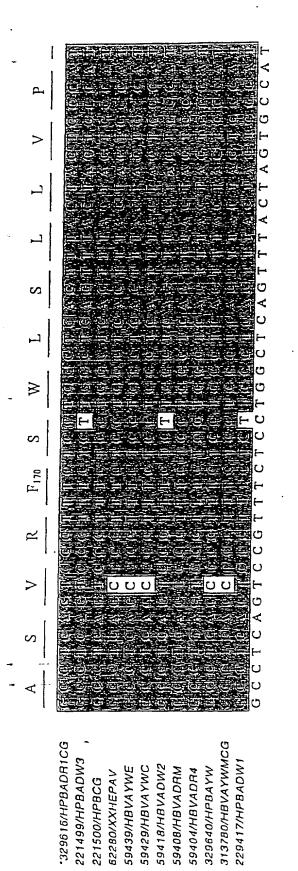
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*329616/HPBADR1CG

221499/HPBADW3

221500/HPBCG



L192 W 191 V_{180}

313780/HBVAYWMCG

329640/HPBAYW

59418/HBVADW2

59408/HBVADRM

59404/HBVADR4

59439/HBVAYWE 59429/HBVAYWC

62280/XXHEPAV

229417/HPBADW1

The High has been done on the same

W. L. M. W. S. J. J. Con. L. J.

Figure 3 continued

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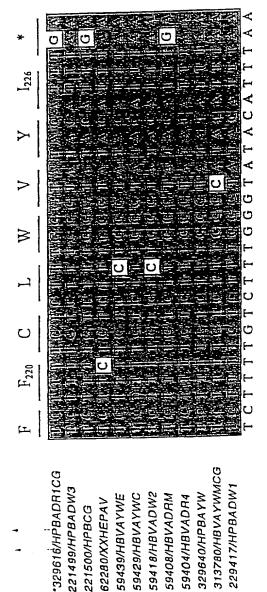


Figure 3 continued

313780/HBVAYWMCG 229417/HPBADW1 329640/HPBAYW 59404/HBVADR4

221499/HPBADW3

59439/HBVAYWE 59429/HBVAYWC 59418/HBVADW2 59408/HBVADAM

62280/XXHEPAV 221500/HPBCG

pBBHBV1.28

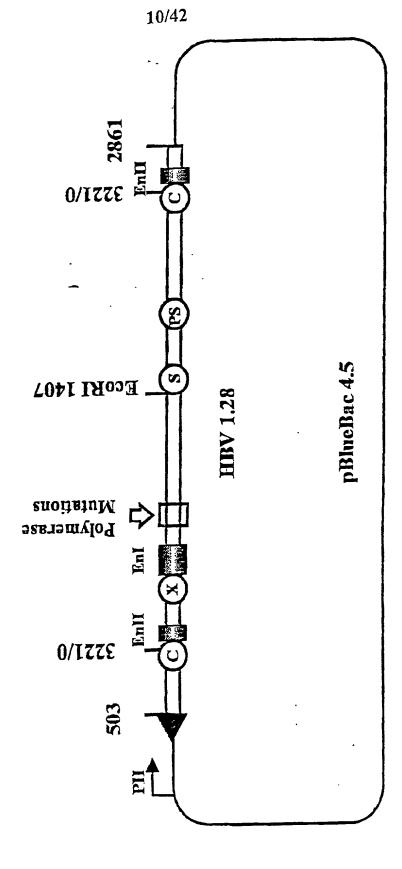


Figure 4A

TE.

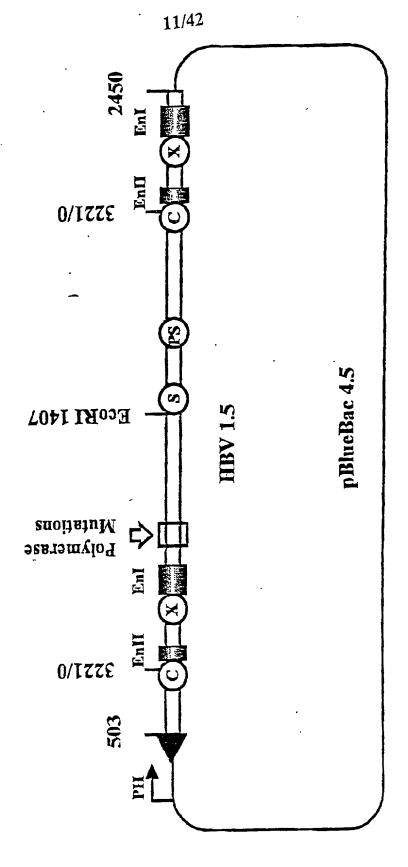


Figure 4B

Sequence Range: 1 to 4084

Figure 5A

Figure 5A continued

	3710	3720	3730	3740	3750	
GCTAC	TGTGGAGTTA	CTCTCGTTT1	TGCCTTCTGA	CTTCTTTCCT	TCCGT	
	•					
	3760	3770	3780	3790	3800	
CAGAG	ATCTCCTAGA	CACCGCCTCA	GCTCTGTATC	GAGAAGCCTT	AGAGT	
	3810		3830	3840	3850	
CTCCT	GAGCATTGCT	CACCTCACCA	TACTGCACTC	AGGCAAGCCA	TTCTC	
	3860	3870		3890	3900	
TGCTG	GGGGGAATTG	ATGACTCTAG	CTACCTGGGT	GGGTAATAAT	ITGGA	
_						
	3910 `		3930	3940	3950	
AGATCCAGCATCCAGGGATCTAGTAGTCAATTATGTTAATACTAACATGG						
	3960	3970	3300	3990	4000	
GTTTAAAGATCAGGCAACTATTGTGGTTTCATATATCTTGCCTTACTTTT						
	4010	4020		4040	4050	
GGAAGAGAGACTGTACTTGAATATTTGGTCTCTTTCGGAGTGTGGATTCG						
	4060		4080	•		
CACTCCTCCAGCCTATAGACCACCAAATGCCCCT						

Figure 5A continued

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7.5

Figure 5B

670 680

CCGACCTTGAGGCCTACTTCAAAGACTGTGTTTTAAGGACTGGGAGGAG

690

700

660

	3710	3720	3730	3740	3750
ACG	GGCGCACC	TCTCTTTACGC	GGTCTCCCC	GTCTGTGCCTT	CTCATCT
		2024		2200	2000
CCC	3760 Saracaran	3770 GCACTTCGCTT	-		3800 3800 CCRC
GCCC	301001010	3CAC11CGC11	CMCCICIGC	ACG11GCA1G0	ENGACCAC
	3810	3820	3830	3840	3850
CGT	AACGCCCA	r CAGAT CCTGC	CCAAGGTCT:	TACATAAGAGG	ACTCTTG
	2060	2070	2000	2000	2000
ርልሮባ	038E CCCAGCAA'	3870 IGTCAACGACC			3900 3GACTGT
- Crac-1	.cccaccar.	re i a precince.	once i i ener	recine i com	ITOFIC I GI
	3910	3920	3930	3940	3950
GTGI	TTAAGGACT	CGGGAGGAGCT	GGGGAGGAG	SATTAGGTTAA	AGGTCTT
	3960	3970	3000	3000	4000
TGTA		3970 TGTAGGCATA			
				4040	
ACTT	TTTCACCTC	TGCCTAATCAT	CTCTTGTAC	ATGTCCCACT	GTTCAAG
	4060	4070	4080	4090	4100
CCTC		CCTTGGGTGGC			
				4140	
AAAG	AATT"IGGAG	CTACTGTGGAG	TTACTCTCG	TTTTTGCCTTC	CTGACTT
	4160	4170	4180	4190	4200
CTTT	CTTCCGTC	AGAGATCTCCT	AGACACCGC	CTCAGCTCTGT	TATCGAG
አአርርር		4220 ICCTGAGCATT	4230		4250
AAGC	LITAGAGIC	ICCIGAGCALI	GCICACCICA	ACCATACTGCA	LL LAGG
	4260	4270	4280	4290	4300
CAAGO	CATTCTCT	GCTGGGGGGAA'	TTGATGACT	CTAGCTACCTG	GGTGGG
	4210	1200	4220		
מית מית	4310	4320 BATCCAGCATC		4340 \CTX CTCX X TT	
TIMITI	MIT TIOOM	JAI COAUCAI C	TUGGGET C IL	GIAGICAAII	AIGIIA
	4360	4370	4380	4390	4400
ATACI	'AACATGGG'	TTAAAGATCA(GCAACTAT1	TGTGGTTTCAT	ATATCT
		4400			
ጥርረረጥ		4420 BAAGAGAGACTO			4450
10001	1WC111100	1 JADADADAN	IMCIIGMAI	.A.11661616	111000
	4460	4470		4490	
AGTG	rggattcgc.	ACTCCTCCAGC	CTATAGACC	ACCAAATGCCC	CT

100

10



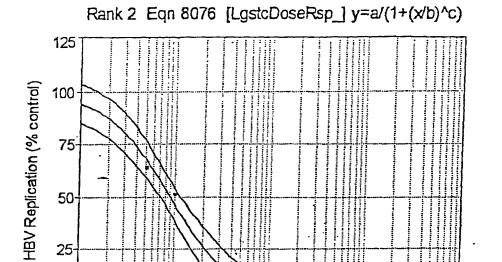


Figure 6A

microMolar 3TC

0.1

0.01

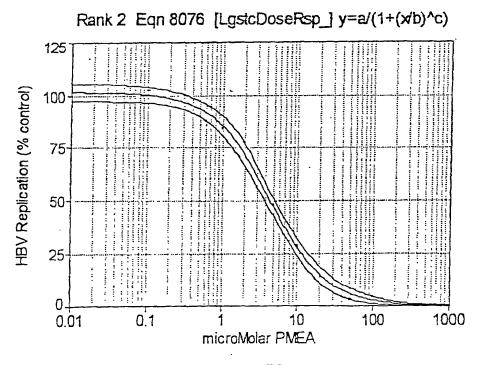


Figure 6B

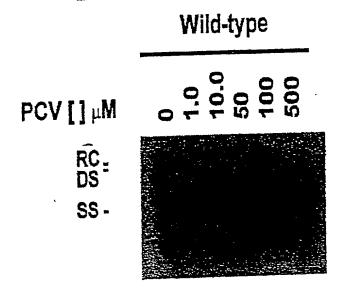


Figure 6C

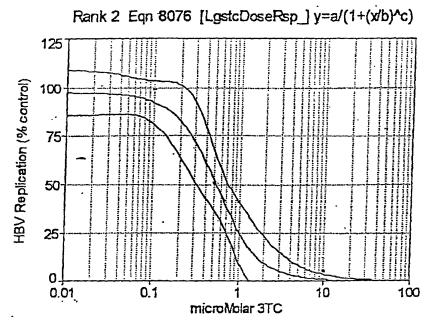


Figure 7A

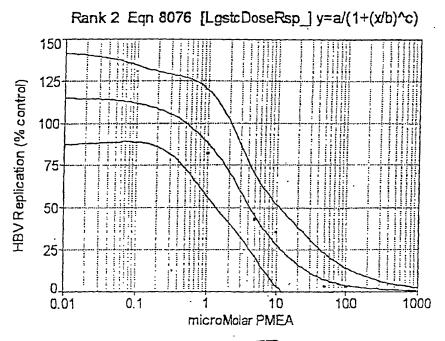


Figure 7B

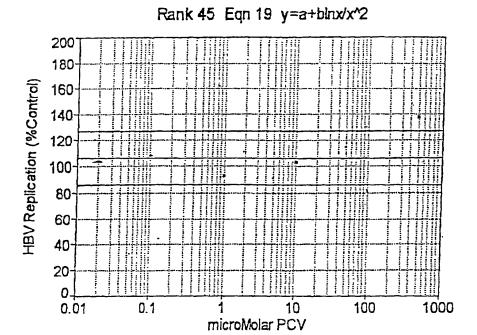
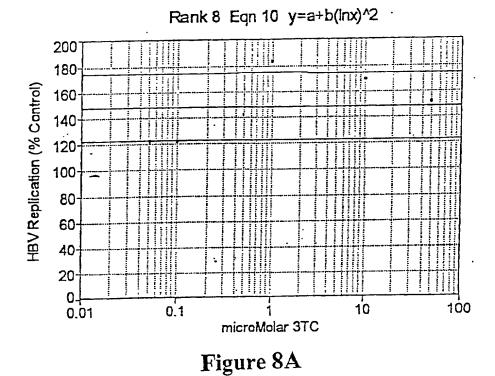


Figure 7C

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Rank 2 Eqn 8076 [LgstcDoseRsp_] y=a/(1+(x/b)^c)

175

150

100

100

75

0

0.01

0.1

100
1000

microMolar PMEA

Figure 8B

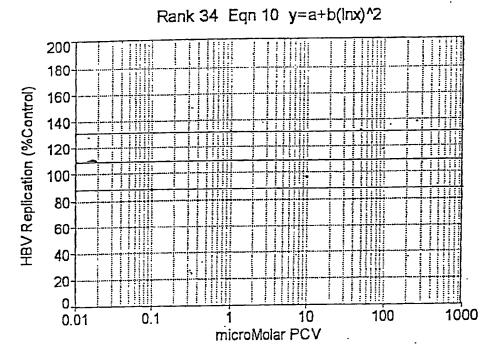


Figure 8C

1



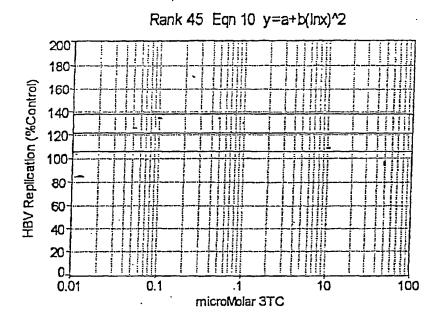


Figure 9A

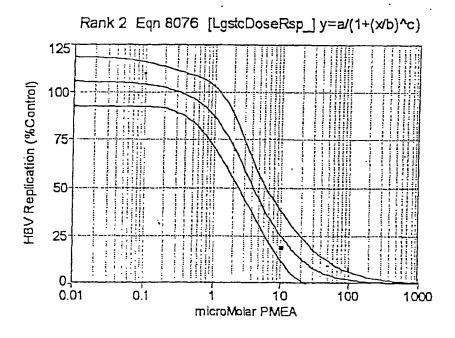


Figure 9B

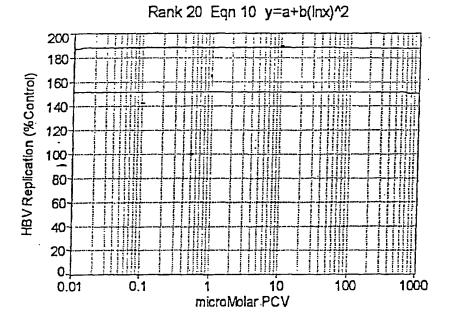


Figure 9C

Cold dCTP Competition

Rank 2 Eqn 8076 [LgstcDoseRsp_j y=a/(1+(x/b)^c)

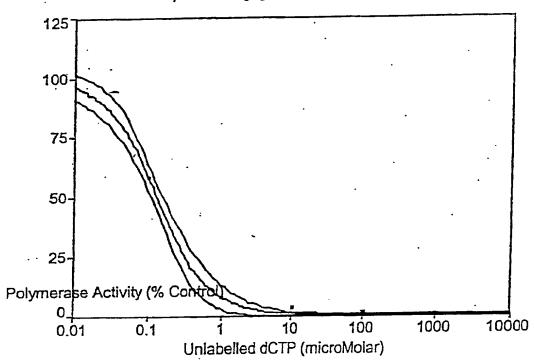


Figure 10

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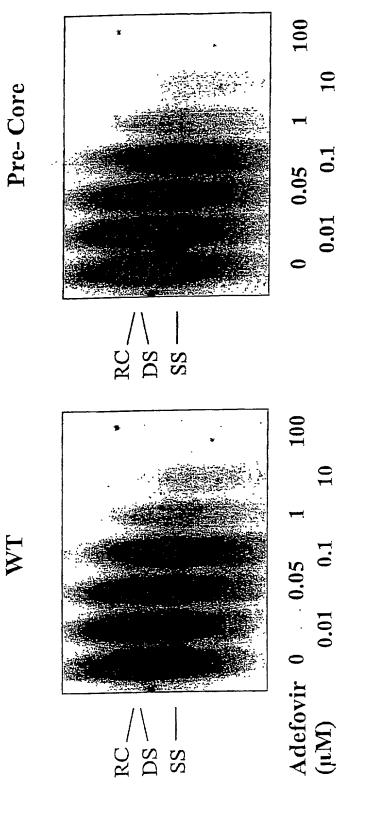


Figure 11A

1

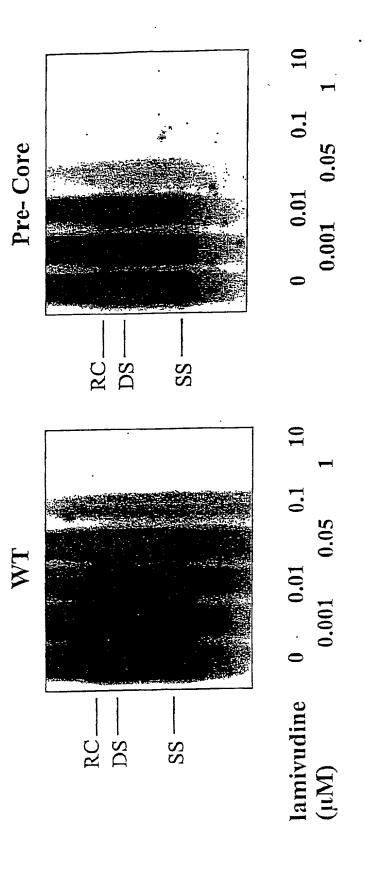


Figure 11B

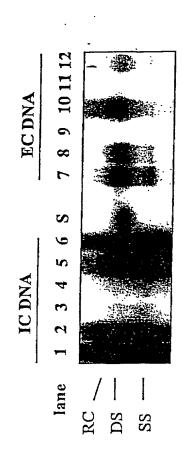
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500

Figure 11C

H .:

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Lane 1 & 7

Lane 2 & 8

Lane 3 & 9

Lane 4 & 11

Lane 5 & 10

Lane 6 & 12

- Wild type (HBV x 1.3) - pre-core - M550I

- Standard

- L526M/M550V

- pre-core/M550I

- pre-core/L526M/M550V

Figure 12

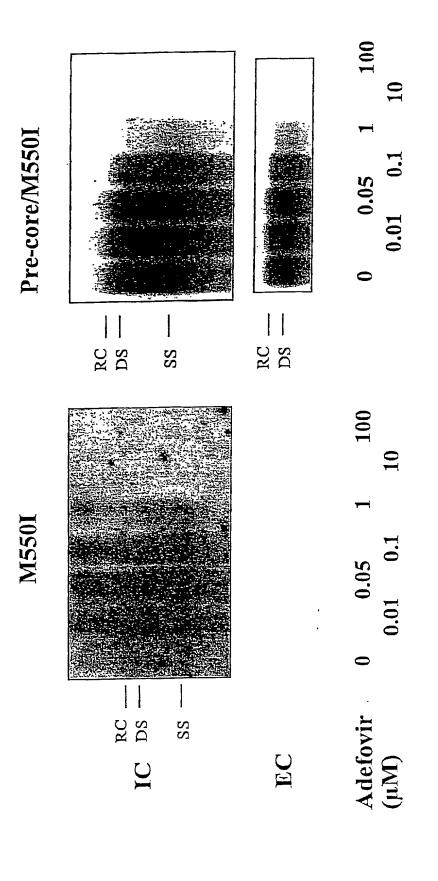


Figure 13A

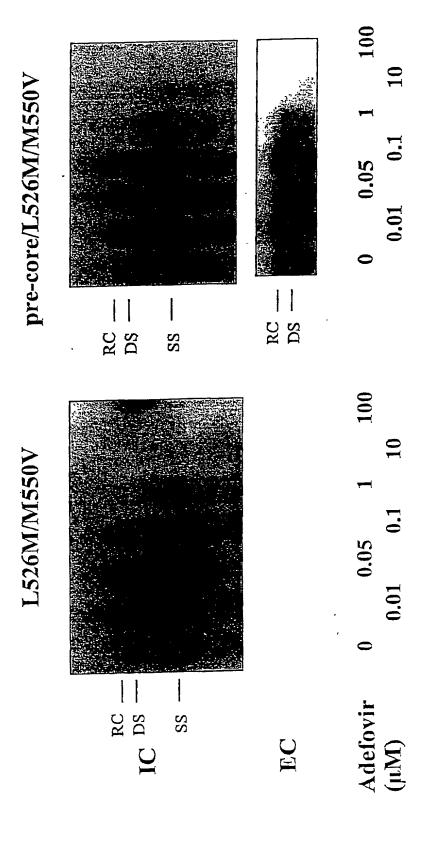


Figure 13B

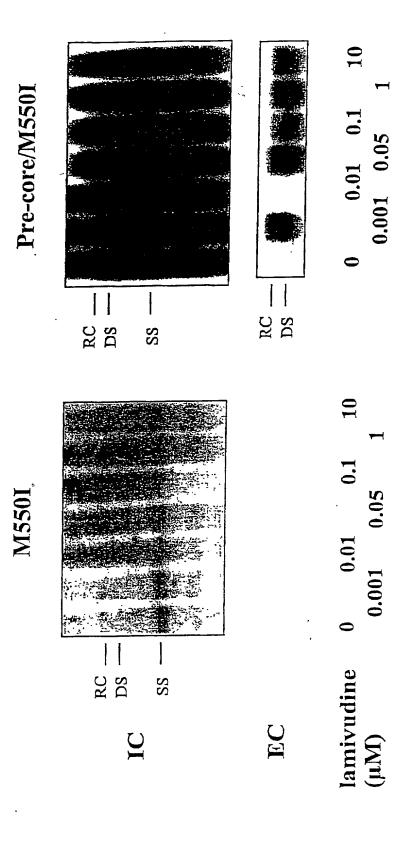


Figure 13C

Figure 13D

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Figure 13E

The state of the s

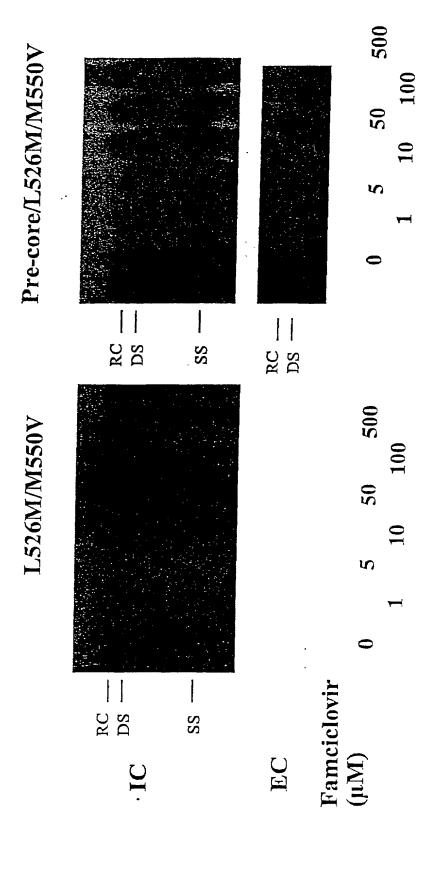


Figure 13F